SEQUENCE LISTING





<110> Innogenetics N.V.

<120> New sequences of hepatitis C virus genotypes for diagnosis, prophylaxis and therapy

- <130> 35
- <160> 284
- <170> PatentIn version 3.1
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- <211> 213
- <212> DNA
- <213> hepatitis C virus
- <400> 1

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- <211> 71
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Ala Gln Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr 20 25 30	
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v

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Ser	Ala	Cys	Trp 100	Thr	Pro	Val	Thr	Pro 105	Thr	Val	Ala	Val	Arg 110	Tyr	Val
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55

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val

75

60

120

180

240

300

360

420

480

540

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Gly	Ala	Thr 115	Thr	Ala	Ser	Ile	Arg 120	Arg	His	Val	Asp	Ile 125	Leu	Val	Gly		
Ala	Ala 130	Thr	Met	Cys	Ser	Ala 135	Leu	Tyr	Val	Gly	Asp 140	Met	Cys	Gly	Ala		
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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Ala Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Arg His Val Asp Ile Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

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His Arg Met Ala 180

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<212> PRT

<213> hepatitis C virus

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val

110 105 100

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 115 Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 150 145 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 170 His Arg Met Ala 180 21 <210> <211> 541 DNA <212> hepatitis C virus <213> <400> 21 cgtcggcgct cccgtaggag gcgtcgcaag agcccttgcg catggcgtga gggcccttga 60 agacgggata aatttcgcaa cagggaattt gcccggttgc tecttttcta ttttccttct 120 tgctctgttc tcttgcttaa ttcatccagc agctagtcta gagtggcgga atacgtctgg 180 cctctatgtc cttaccaacg actgttccaa tagcagtatt gtgtacgagg ccgatgacgt 240 tattctgcac acacccggct gcataccttg tgtccaggac ggcaatacat ccacgtgctg 300 gaccccagtg acacctacag tggcagtcaa gtacgtcgga gcaaccaccg cttcgatacg 360 cagtcatgtg gacctattag tgggcgcggc cacgatgtgc tctgcgctct acgtgggtga 420 catgtgtggg gctgtcttcc tcgtgggaca agccttcacg ttcagacctc gtcgccatca 480 aacggtccag acctgtaact gctcgctgta cccaggccat ctttcaggac atcgaatggc 540 541 t <210> 22 <211> 180

PRT <212>

<213> hepatitis C virus

<400> 22

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 25

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 55

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 90 85

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val 110 105 100

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 120 115

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 150 145

Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 170 165

His Arg Met Ala 180

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85. 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val
100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 150 160

Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 165 170 175

His Arg Met Ala 180

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 45 35 Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 75 70 Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val 105 100 Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 125 120 115 Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 150 145 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 175 170 165 His Arg Met Ala 180 <210> 27 <211> 541 <212> DNA <213> hepatitis C virus <400> 27 cgtcggcgct cccgtaggag gcgtcgcaag agcccttgcg catggcgtga gggcccttga 60 ggacgggata aacttcgcaa cagggaattt gcccggttgc tctttttcta tcttccttct 120 tgctctgttc tcttgcttaa tccatccagc agctggtcta gagtggcgga atacgtctgg 180 cctctatgtc cttaccaacg actgttccaa tagtagtatt gtgtatgagg ccgatgacgt 240 tattctgcac gcgcccggct gtgtaccttg tgtccaggac ggcaatacgt ctacatgctg 300 gaccccagta acacctacag tggcagtcag gtacgtcggg gcaaccaccg cttcgatacg 360
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<212> PRT

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<400> 28

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 165 His Arg Met Ala 180 <210> 29 <211> 287 <212> DNA <213> hepatitis C virus <400> 29 tagacttttg ggagagcgtc ttcactggac taactcacat agatgcccac tttctgtcac 60 agactaagca gcagggactc aacttctcgt tcctgactgc ctaccaagcc actgtgtgcg 120 180 ctcgcgcgca ggctcctccc ccaagttggg acgagatgtg gaagtgtctc gtacggctta agccaacact acatggacct acgcctcttc tatatcggtt ggggcctgtc caaaatgaaa 240 287 totgottgac acaccocato acaaaataca toatggcatg catgtca <210> 30 <211> 95 <212> PRT <213> hepatitis C virus <400> 30 Asp Phe Trp Glu Ser Val Phe Thr Gly Leu Thr His Ile Asp Ala His 5 Phe Leu Ser Gln Thr Lys Gln Gln Gly Leu Asn Phe Ser Phe Leu Thr 25 20 Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser 35 Trp Asp Glu Met Trp Lys Cys Leu Val Arg Leu Lys Pro Thr Leu His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 75 70

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser

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Cys	Met	Ser	Ala 20	Asp	Leu	Glu	Val	Thr 25	Thr	Ser	Thr	Trp	Val 30	Leu	Leu
Gly	Gly	Val 35	Leu	Ala	Ala	Leu	Ala 40	Ala	Tyr	Cys	Leu	Ser 45	Val	Gly	Cys
Val	Val 50	Ile	Val	Gly	His	Ile 55	Glu	Leu	Gly	Gly	Lys 60	Pro	Ala	Leu	Val
Pro 65	Asp	Lys	Glu	Val	Leu 70	Tyr	Gln	Gln	Tyr	Asp 75	Glu	Met	Glu	Glu	Cys 80
Ser	Gln	. Ala	Ala	Pro 85	Tyr	Ile	Glu	Gln	Ala 90	Gln	Val	Ile	Ala	His 95	Gln
Phe	Lys	Glu	Lys 100	Ile	Leu	Gly	Leu	Leu 105	Gln	Arg	Ala	Thr	Gln 110	Gln	Gln

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr

125

Phe Trp His Lys His

<210> 33 <211> 401 <212> DNA <213> hepatitis C virus <400> 33 tocaaaatga aatotgottg acacaccocg toacaaaata cattatggca tgcatgtcag 60 ctgatctgga agtaaccacc agcacctggg tgttgcttgg aggggtcctc gcggccctag 120 cggcctactg cttgtcagtc ggctgcgttg taatcgtggg tcatatcgag ctggggggca 180 240 agccqqcact cgttccagac aaggaggtgt tgtatcaaca gtacgatgag atggaggagt qctcgcaagc cgccccatac atcgaacaag ctcaggtaat agcccaccag ttcaaggaga 300 aaatccttgg actgctgcag cgagccaccc aacaacaagc tgtcattgag cccgtaatag 360 401 cttccaactg gcaaaagctt gaaaccttct ggcacaagca t <210> 34

<211> 133

<212> PRT

<213> hepatitis C virus

<400> 34

Gln Asn Glu Ile Cys Leu Thr His Pro Val Thr Lys Tyr Ile Met Ala 1 5 10 15

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Leu Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 65 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Ile Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln 100 105 110

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr 115 120 125

<210> 35 <211> 401 <212> DNA <213> hepatitis C virus

130

<400> 35
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ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120
cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca 180
agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240
gctcacaagc tgccccatat atcgaacaag ctcaggtaat agctcaccag ttcaagggaa 300
aagtccttgg attgctgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360
ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t

<210> 36 <211> 133 <212> PRT <213> hepatitis C virus

<400> 36

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala 1 5 10 15

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Gly Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 105 100 Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 37 <211> 401 DNA <212> <213> hepatitis C virus <400> 37 tccaaaatga aatctgcttg acacacccca tcacaaaata catcatggca tgcatgtcag 60 ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 eggeetactg ettgteagte ggttgtgttg tgattgtggg teatategag etggggggea 180 agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 gctcacaagc tgccccatat atcgaacaag ctcaggtgat agctcaccag ttcaaggaaa 300 aagtccttgg attgctgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360 401 ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t <210> 38 <211> 133 PRT <212> <213> hepatitis C virus <400> .38 Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 25

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys
35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val
50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

75

Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln 100 105 110

Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 115 120 125

Phe Trp His Lys His 130

<210> 39 <211> 401 <212> DNA <213> hepatitis C virus

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<400> 39
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ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120
cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca 180
agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240
gctcacaagc tgccccatat atcgaacaag ctcaggtaat agctcaccag ttcaaggaaa 300
aagtccttgg attgctgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360
ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t 401

<210> 40 <211> 133 <212> PRT <213> hepatitis C virus

<400> 40

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

5

Cvs	Met	Ser	Ala	Asp	Leu	Glu	Val	Thr	Thr	Ser	Thr	Trp	Val	Leu	Lėu
Cy S			20	•				25					30		

10

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 65 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln 100 105 110

Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 115 120 125.

Phe Trp His Lys His

<210> 41

<211> 509

<212> DNA

<213> hepatitis C virus

<400> 41 ccatgagcac gaatcctaaa cctcaaagaa aaaccaaaag aaacaccaac cgtcgcccac 60 aggacgtcaa gttcccgggc ggtggtcaga tcgttggcgg agtttacttg ttgccgcgca 120 ggggccctag gatgggtgtg cgcgcgactc ggaagacttc ggaacggtcg caaccccgtg 180 gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 240 ggtaccettg geceetttae gecaatgagg geetegggtg ggeagggtgg etgeteteee 300 ctcgaggetc tcggcctaat tggggcccca atgacccccg gcgaaaatcg cgtaatttgg 360 gtaaggtcat cgatacccta acgtgcggat tcgccgatct catggggtat atcccgctcg 420 taggcggccc cattgggggc gtcgcaaggg ctctcgcaca cggtgtgagg gtccttgagg 480 <210> 42

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 42

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 43

<211> 509

<212> DNA

<213> hepatitis C virus

<400> 43			•			
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ggggccctag	gatgggtgtg	cgcgcgactc	ggaagacttc	ggaacggtcg	caaccccgtg	180
gacggcgtca	gcctattccc	aaggcgcgcc	agcccacggg	ccggtcctgg	ggtcaacccg	240
ggtacccttg	gcccctttac	gccaatgagg	gcctcgggtg	ggcagggtgg	ctgctctccc	300
ctcgaggctc	tcggcctaat	tggggcccca	atgacccccg	gcgaaaatcg	cgtaatttgg	360
gtaaggtcat	cgatacccta	acgtgcggat	tcgccgatct	catggggtat	atcccgctcg	420
taggcggccc	cattgggggc	gtcgcaaggg	ctctcgcaca	cggtgtgagg	gtccttgagg	480
acggggtaaa	ctatgcaaca	gggaattta				509

<210> 44

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 44

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp . 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 45

<211> 580

<212> DNA

<213> hepatitis C virus

<400> 45

aacgtgcgga ttcgccgatc tcatggggta tatcccgctc gtaggcggcc ccattggggg 60 cgtcgcaagg gctctcgcac acggtgtgag ggtccttgag gacggggtaa actatgcaac 120 agggaattta cccggttgct ctttctctat ctttattctt gctcttctct cgtgtctgac 180 cqttccggcc tctgcagttc cctaccgaaa tgcctctggg atttatcatg ttaccaatga 240 300 ttqcccaaac tcttccataq tctatqaggc agataacctg atcctacacg cacctggttg 360 cqtqccttqt qtcatgacag gtaatgtgag tagatgctgg gtccaaatta cccctacact 420 qtcaqccccg agcctcggag cagtcacggc tcctcttcgg agagccgttg actacctagc gggagggct gccctctgct ccgcgttata cgtaggagac gcgtgtgggg cactattctt 480 ggtaggccaa atgttcacct ataggcctcg ccagcacgct acggtgcaga actgcaactg 540 580 ttccatttac agtggccatg ttaccggcca ccggatggca

<210> 46

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 46

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

الم الم

Glu	Asp	Gly 35	Val	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Сув	Ser	Phe	
Ser	Ile 50	Phe	Ile	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Ala	Ser	
Ala 65	Val	Pro	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80	
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	Asn	Leu	Ile	Leu 95	His	
 Ala	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Met 105	Thr	Gly	Asn	Val	Ser 110	Arg	Cys	
Trp	Val	Gln 115	Ile	Thr	Pro	Thr	Leu 120	Ser	Ala	Pro	Ser	Leu 125	Gly	Ala	Val	•
Thr	Ala 130	Pro	Leu	Arg	Arg	Ala 135	Val	Asp	Tyr	Leu	Ala 140	Gly	Gly	Ala	Ala	
Leu 145	Cys	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Ala	Cys 155	Gly	Ala	Leu	Phe	Leu 160	
Val	Gly	Gln	Met	Phe 165	Thr	Tyr	Arg	Pro	Arg 170	Gln	His	Ala	Thr	Val 175	Gln	
Asn	Cys	Asn	Cys 180	Ser	Ile	Tyr	Ser	Gly 185	His	Val	Thr	Gly	His 190	Arg	Met	
Ala																
<210 <210 <210 <210	1> 5 2> I	17 580 DNA nepat	:itis	3 C 1	/irus	5										
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															gcaac	12
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cgttccggcc tctgcagttc cctaccgaaa tgcctctggg atttatcatg ttaccaatga 240
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cgtgccttgt gtcatgacag gtaatgtgag tagatgctgg gtccaaatta cccctacact 360
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gggaggggct gccctctgct ccgcgttata cgtaggagac gcgtgtgggg cactattctt 480
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<210> 48

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 48

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys 100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135 140 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 175

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210> 49

<211> 959

<212> DNA

<213> hepatitis C virus

<400> 49 ccatgagcac gaatcctaaa cctcaaagaa aaaccaaaag aaacaccaac cgtcgcccac 60 aggacgtcaa gttcccgggc ggtggtcaga tcgttggcgg agtttacttg ttgccgcgca 120 ggggccctag gatgggtgtg cgcgcgactc ggaagacttc ggaacggtcg caaccccgtg 180 gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 240 300 ggtaccettg geceetttae gecaatgagg geetegggtg ggeagggtgg etgeteteee 360 ctcgaggete teggeetaat tggggeecea atgaceeeg gegaaaateg egtaatttgg gtaaggtcat cgatacccta acgtgcggat tcgccgatct catggggtat atcccgctcg 420 480 taggcggccc cattgggggc gtcgcaaggg ctctcgcaca cggtgtgagg gtccttgagg acggggtaaa ctatgcaaca gggaatttac ccggttgctc tttctctatc tttattcttg 540 600 ctcttctctc gtgtctgacc gttccggcct ctgcagttcc ctaccgaaat gcctctggga 660 tttatcatgt taccaatgat tgcccaaact cttccatagt ctatgaggca gataacctga tectacaege acctggttge gtgeettgtg teatgacagg taatgtgagt agatgetggg 720 tecaaattae ecctacaetg teageceega geeteggage agteaegget eetettegga 780 gageegttga etaeetageg ggaggggetg ceetetgete egegttatae gtaggagaeg 840 900 cgtgtggggc actattcttg gtaggccaaa tgttcaccta taggcctcgc cagcacgcta 959 cggtgcagaa ctgcaactgt tccatttaca gtggccatgt taccggccac cggatggca

<211> 319

<212> PRT

<213> hepatitis C virus

<400> 50

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 . 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 290 295 300

Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 305 310 315

<210> 51

<211> 959

<212> DNA

<213> hepatitis C virus

<400> 51

ccatgagcac gaatcctaaa cctcaaagaa aaaccaaaag aaacaccaac cgtcgcccac 60 aggacqtcaa gttcccgggc ggtggtcaga tcgttggcgg agtttacttg ttgccgcgca 120 ggggccctag gatgggtgtg cgcgcgactc ggaagacttc ggaacggtcg caaccccgtg 180 gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 240 300 ggtaccettg gcccctttac gccaatgagg gcctcgggtg ggcagggtgg ctgctctccc ctcgaqqctc tcggcctaat tggggcccca atgacccccg gcgaaaatcg cgtaatttgg 360 qtaaqqtcat cgatacccta acgtgcggat tcgccgatct catggggtac atcccgctcg 420 taggcqqccc cgttgggggc gtcgcaaggg ctctcgcaca cggtgtgagg gtccttgagg 480 acggggtaaa ctatccaaca gggaatttac ccggttgctc tttctctatc tttattcttg 540 ctcttctctc gtgtctgacc gttccggcct ctgcagttcc ctaccgaaat gcctctggga 600 660 tttatcatgt taccaatgat tgcccaaact cttccatagt ctatgaggca gataacctga

tectacaege acetegetes greectege teatgacaege taatgrage agategeggg 720
tecaaattae ecetacaeteg teageecega geeteggage agteaegget cetettegga 780
gageegttga etacetagee ggaggggeteg ecetetegete egegttatae graggagaeeg 840
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<210> 52

<211> 319

<212> PRT

<213> hepatitis C virus

<400> 52

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 170 165 Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 185 180 Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 200 195 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 215 210 Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 235 230 225 Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 280 Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 300 295 290 Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 310 305 <210> 53 <211> 959 <212> DNA <213> hepatitis C virus <400> 53 ccatgagcac gaatcctaaa cctcaaagaa aaaccaaaag aaacaccaac cgtcgcccac 60 aggacgtcaa gttcccgggc ggtggtcaga tcgttggcgg agtttacttg ttgccgcgca 120 ggggccctag gatgggtgtg cgcgcgactc ggaagacttc ggaacggtcg caaccccgtg 180 gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 240 300 ggtaccettg gcccetttac gccaatgagg gcctcgggtg ggcagggtgg ctgctctccc

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<210> 54

<211> 319

<212> PRT

<213> hepatitis C virus

<400> 54

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 · 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205

Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

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120

180

240

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Lys Asn Cys Asp Glu Leu 115

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attcttggga taggcactgt ccttgaccag gcagagacgg ctggagctag gctcgtcgtc 180
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Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr Val Leu
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Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Xaa Thr Xaa
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Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val
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65
Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Xaa Arg Ala Ile Pro
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Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys
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105

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Arg Pro Trp Glu Tyr 130

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atcatcctct ctgggaaacc tgccatcatt cccgataggg aggcattata ccancaattt
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                                                                     240
gatgagatgg aggagtgctc ggcctcgttg ccctatatgg acgagacacg tgccattgcc
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qqacaattca aagagaaagt gctcggcttc atcagcacga ccggccagaa ggctgaaact
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Ile Ile Pro Asp Arg Glu Ala Leu Tyr Xaa Gln Phe Asp Glu Met Glu 55 50

Glu Cys Ser Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala 75 70 65

Gly Gln Phe Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln 95

Lys Ala Glu Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala 105 100

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	3> artificial	
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cut		
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Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Lys Ala 65 70 75 80

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20 25 30

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35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

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Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
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Ala

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Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ser Gly Leu Arg Asp Cys Thr Met Leu Val Tyr Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
100 105 110

Val

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caccaccagc ttcgggaaca cactgacgtg ctacctcaaa gcctcagccg ctatcagagc 240
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Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
100 105 110

Ala

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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr 20 25 30											
Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45											
Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60											
Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Val Ala Ala Thr Arg Ala 65 70 75 80											
Ala Gly Leu Lys Gly Cys Ser Met Leu Val Cys Gly Asp Asp Leu Val											

Val Ile Cys Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg

Ala

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gga	gccc	gag	gccc	gcaa	gg t	aata	gccg	c cc	tcac	ggag	aga	ctct	acg	tggg	cggccc
cat	gcat	aac	agca	aggg	ag a	cctt	tgcg	g gt	atcg	taga	tgc	cgcg	cga	gcgg	cgtata
cac	cacc	agc	ttcg	ggaa	ca c	aatg	acgt	g ct	acct	taag	gcc	tcag	cag	ccat	cagggc
tgc	3999	cta	aagg	attg	ca c	catg	ctgg	t tt	gcgg	tgac	gac	ctag	tcg	tgat	cgccga
gag	ggt	ggc	gttġ	agga	gg a	caaa	cgag	c cc	tcgg	agct					
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Ser 1	Thr	Val	Thr	Glu 5	Arg	Asp	Ile	Lys	Val 10	Glu	Glu	Glu	Val	Tyr 15	Gln
Cys	Cys	Asp	Leu 20	Glu	Pro	Glu	Ala	Arg 25	Lys	Val	Ile	Ala	Ala 30	Leu	Thr
Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	His	Asn	Ser	Lys 45	Gly	Asp	.Leu
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala		Gly	Val	Tyr 60	Thr	Thr	Ser	Phe
Gly 65	Asn	Thr	Met	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Ile	Arg	Ala 80
Ala	Gly	Leu	Lys	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val
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Ala

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Pro Val Gly Gly Val Ala Arg	Ala Leu Ala His 25	Gly Val Arg Ala Val 30
Glu Asp Gly Ile Asn Tyr Ala 35	Thr Gly Asn Leu 40	Pro Gly Cys Ser Phe 45
Ser Ile Phe Leu Leu Ala Leu 50 55		Thr Val Pro Thr Ser
Ala Val Asn Tyr Arg Asn Ala 65 70	Ser Gly Ile Tyr 75	His Ile Thr Asn Asp 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His

Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 120 125

Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg 180 185 190

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<213> hepatitis C virus

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<210> 121

<211> 191

<212> PRT

<213> hepatitis C virus

<400> 121

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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys
100 105 110

Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Val Gly Ala Pro
115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg 180 185 190

<210> 122

<211> 574

<212> DNA

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<400> 122

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gggaatcttc	ccggttgctc	tttctctatc	ttcctcctgg	cacttctttc	gtgcctcact	180
gtcccagcgt	cagctgagca	ctaccggaat	gcttcgggca	tctatcacat	caccaatgac	240
tgtccgaatt	ccagcgtagt	ctatgaaact	gaccaccata	tattgcactt	gccggggtgc	300
gtaccctgcg	tgagggccgg	gaacgtgtct	cgttgctgga	cgccggtaac	acctacggtg	360
gctgccgtat	ccatggacgc	tccgctcgag	teetteegge	ggcatgtgga	cctaatggta	420
ggtgcggcca	ccgtgtgttc	tgtcctctat	gttggagacc	tctgtggagg	tgctttccta	480
gtggggcaga	tgttcacctt	ccagccgcgt	cgccactgga	ccacgcagga	ttgtaattgc	540
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<210> 123

<211> 191

<212> PRT

<213> hepatitis C virus

<400> 123

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Val Val Tyr Glu Thr Asp His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Ala Gly Asn Val Ser Arg Cys

Trp Thr Pro Val Thr Pro Thr Val Ala Ala Val Ser Met Asp Ala Pro 115 120 125

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Leu Glu Ser Phe Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
    130
                        135
Val Cys Ser Val Leu Tyr Val Gly Asp Leu Cys Gly Gly Ala Phe Leu
                    150
                                        155
145
Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln
                                    170
                165
Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg
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Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val
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Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile
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Val Tyr Glu Thr Glu His His Ile Leu His Leu
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Val Tyr Glu Ala Asp His His Ile Met His Leu
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Val Tyr Glu Thr Asp His His Ile Leu His Leu
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Val Arg Ala Gly Asn Val Ser Arg Cys Trp Thr Pro Val
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Ala Pro Tyr Ile Gly Ala Pro Leu Glu Ser
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Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser
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Gln Pro Arg Arg His Trp Thr Thr Gln Asp
               5
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Arg Pro Arg Arg His Trp Thr Thr Gln Asp
             5
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Arg Pro Arg Arg His Trp Thr Thr Gln Asp
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Asp Arg Arg Pro Thr Gly Lys Ser Trp Gly Lys Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp
85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro
100 105 110

Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Val Pro Val Ser Gly Leu 180 185 190

Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Gln 195 200 205

Asn Ser Ser Ile Val Trp Gln Leu Arg Asp Ala Val Leu His Val Pro 210 215 220

Gly Cys Val Pro Cys Glu Glu Lys Gly Asn Ile Ser Arg Cys Trp Ile

Pro Val Ser Pro Asn Ile Ala Val Ser Gln Pro Gly Ala Leu Thr Lys 245 250 255

Gly Leu Arg Thr His Ile Asp Thr Ile Ile Ala Ser Ala Thr Phe Cys 260 265 270

Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Ala Val Met Leu Ala Ser 275 280 285

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225

<211> 340

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<213> hepatitis C virus

<400> 145

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caccactagc atgggaaata ccatcacgtg ctacgtgaag gcacaagcag cctgtaaggc 240
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gagtcagggg accgaggagg acgagcggaa cctacgagcc 340

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<211> 113

<212> PRT

<213> hepatitis C virus

<400> 146

Ser Thr Val Thr Glu Arg Asp Ile Arg Thr Glu Glu Ser Ile Tyr Leu
1 5 10 15

Ala Cys Ser Leu Pro Glu Gln Ala Arg Thr Ala Ile His Ser Leu Thr 20 25 30

Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Leu	Asn	Ser	Lys 45	Gly	Gln	Thr	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	
Gly 65	Asn	Thr	Ile	Thr	Cys 70	Tyr	Val	Lys	Ala	Gln 75	Ala	Ala	Суз	Lys	Ala 80	
Ala	Gly	Ile	Ile	Ala 85	Pro	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Ser	Glu 100	Ser	Gln	Gly	Thr	Glu 105	Glu	Asp	Glu	Arg	Asn 110	Leu	Arg	
Ala																
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	_	_													cagta	180
									•						egggt	240
					_								tc c	tgtc	cccgc	300
gcgg	ctct	.cg c	ccgt	cgtg	19 99	ccca	aatg	acc	cccg	gcg	cagg	a				345
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Met 1	Ser	Thr	Leu	Pro 5	Lys	Pro	Gln	Arg	Lys 10	Thr	Lys	Arg		Thr 15	Asn	
Pro	Gly	His	Arg 20	Thr	Leu	Ser		Gln 25	Ala	Ala	Val		Ser 30	Leu	Val	

Glu Phe Thr Cys Tyr His Ala Gly Ala Pro Ser Trp Val Cys Val Gln 40 Cys Ala Arg Leu Pro Ser Gly Arg Asn Leu Ala Val Gly Ala Asn Pro 55 Ser Pro Gly Arg Ala Glu Pro Arg Ala Gly Pro Gly Leu Ser Pro Gly 70 75 Thr Leu Gly Pro Tyr Met Gly Met Arg Ala Ala Gly Gly Gln Gly Gly 90 Ser Cys Pro Arg Ala Ala Leu Ala Arg Arg Gly Ala Gln Met Thr Pro 100 Gly Ala Gly 115 <210> 149 <211> 280 <212> DNA <213> hepatitis C virus <400> 149 60 qqcctqtgac ctcaaggacg aggctaggag ggtgataact tcactcacgg agcggcttta ctgtggtggt cctatgttca acagcaaggg acaacactgc ggttaccgcc gctgccgtgc 120 tagtggggtg ctacccacca gcttcgggaa cacaatcacc tgttacatca aagcaaaggc 180 agctaccaaa gctgccggaa ttaaaaatcc atcattcctt gtctgcggag atgacttggt 240 280 cgtgattgct gagagtgcag ggatcgatga ggacagagcg <210> 150 <211> 93 <212> PRT <213> hepatitis C virus <400> 150 Ala Cys Asp Leu Lys Asp Glu Ala Arg Arg Val Ile Thr Ser Leu Thr

Glu Arg Leu Tyr Cys Gly Gly Pro Met Phe Asn Ser Lys Gly Gln His

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr Ser Phe 40 35 Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Lys Ala Ala Thr Lys Ala 50 Ala Gly Ile Lys Asn Pro Ser Phe Leu Val Cys Gly Asp Asp Leu Val Val Ile Ala Glu Ser Ala Gly Ile Asp Glu Asp Arg Ala <210> 151 <211> 499 <212> DNA <213> hepatitis C virus <400> 151 atgagcacga atcctaaacc tcaaagaaaa accaaaagaa acaccaaccg tcgcccacag 60 gacgtcaagt tocogggogg tggtcagatc gttggoggag tttacttgtt gccgogcagg 120 180 qqccctagga tgggtgtgcg cgcgactcgg aagacttcgg aacggtcgca accccgtgga 240 cqqcqtcaqc ctattcccaa ggcgcgccag cccacgggcc ggtcctgggg tcaacccggg 300 taccettqqc ccctttacgc caatgagggc ctcgggtggg cagggtggct gctctcccct cqaqqctctc ggcctaattg gggccccaat gacccccggc gaaaatcgcg taatttgggt 360 420 aaggtcatcg ataccctaac gtgcggattc gccgatctca tggggtatat cccgctcgta qqcqqcccca ttgggggcgt cgcaagggct ctcgcacacg gtgtgagggt ccttgaggac 480 499 ggggtaaact atgcaacag <210> 152 <211> 166 <212> PRT

<213> hepatitis C virus

<400> 152

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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala

40 45

35

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 70 75 Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110 Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 120 125 115 Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 150 145 Gly Val Asn Tyr Ala Thr 165 <210> 153 <211> 579 <212> DNA <213> hepatitis C virus <400> 153 60 acgtgcggat tcgccgatct catggggtac atcccgctcg taggcggccc cgttgggggc qtcqcaaqqq ctctcgcaca cggtgtgagg gtccttgagg acggggtaaa ctatccaaca 120 180 qqqaatttac ccqqttqctc tttctctatc tttattcttg ctcttctctc gtgtctgacc qttccqqcct ctqcagttcc ctaccgaaat gcctctggga tttatcatgt taccaatgat 240 tgcccaaact cttccatagt ctatgaggca gataacctga tcctacacgc acctggttgc 300 360 qtqccttqtq tcatqacagg taatgtgagt agatgctggg tccaaattac ccctacactg tcagccccga gcctcggagc agtcacggct cctcttcgga gagccgttga ctacctagcg 420

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<210> 154

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 154

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120 125

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135 140

Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 175

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met

180 185 190

Ala

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gtcgcaa	999	ctctcgcaca	tggtgtgagg	gttcttgagg	acggggtgaa	ttatgcaaca	120
gggaatc	tgc	ctggttgctc	tttctctatc	ttcattcttg	cacttctctc	gtgcctcact	180
gtcccgg	cct	ctgcagttcc	ctaccgaaat	gcctctggga	tctatcatgt	caccaatgat	240
tgcccaa	act	cttccatagt	ctatgaggca	gatgatctga	tcctacacgc	acctggctgc	300
gtgcctt	gtg	tcaggaaaga	taatgtgagt	aggtgctggg	tccaaattac	ccccacgctg	360
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<211> 193

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<400> 156

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Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn 75	3
65 Arg Asn Ala Ser Gly Ile The	
70 His Val Thr Asr	λ
75	av Asb
Cys Pro Asn Ser Ser Ilo W.	
Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu 85	
90 Asp Leu Ile Leu	His
Ala Dec = 2	
Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg	
100 Arg Lys Asp Asp Val C	
105 var Ser Arg	Cys
Trp Val Cln Tl	
Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala V	
120 Ala Pro Ser Phe Gly Ala a	
125 Ala V	al
Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Al 130 135 140	
130 Arg Arg Ala Val Asp Tyr Law 200	
135 Val Gly Gly Ala Al	2
140	·a
Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150	
150 Val Gly Asp Ala Cvs Clarat	
155 Ala Leu Phe Leu	1
Val clus as)
of Gin Met Phe Thr Tyr Are a	
165 Pro Arg Gln His Ala Thu sa	
Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 170	
Asp Cys Asp Cys as	
180 Ser Ile Tyr Ser Gly William	•
180 Tyr Ser Gly His Val Thr Gly His Glr Mat	•
Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 180 185 190	
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<213> hepatitis C virus

<400> 158

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Ala Tyr Tyr Ala Ser Val Ala Asn Trp Thr Lys Val Val Leu Val Leu

Phe Leu Phe Ala Gly Val Asp Ala Thr Thr Gln Ile Ser Gly Gly Ser

Ser Ala Gln Thr Thr Tyr Gly Ile Ala Ser Phe Ile Thr Arg Gly Ala

Gln Gln Lys Leu Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn

Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Ile Ala

Gly Leu Phe Tyr Tyr His Lys Phe Asn Ser Ser Gly Cys Pro Asp Arg

Met Ala Ser Cys Arg Ala Leu Ala Thr Phe Asp Gln Gly Trp Gly Thr

Ile Ser Tyr Ala Asn Ile Ser Gly Pro Ser Asp Asp Lys Pro Tyr Cys

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<211> 113

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Gln Arg Leu Tyr Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln
35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Ala 65 70 75 80

Ala Arg Leu Arg Asp Cys Thr Leu Leu Val Cys Gly Asp Asp Leu Val 85 90 95

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Ala .

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Ser Cys Asp Ser Gln Pro Asp Ala Arg Ala Ala Ile Arg Ser Leu Thr 20 25 30

Gln Arg Leu Phe Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Thr 65 70 75 80

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Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Asn Leu Arg 100 105 110

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360

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480

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85

90

120

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala 130 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 150 145 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 180 Ala <210> 197 <211> 1485 <212> DNA <213> hepatitis C virus <220> <221> misc_feature <222> (787)..(787) <223> "n" is any nucleotide <220> <221> misc_feature <222> (1011)..(1011) <223> "n" is any nucleotide <220> <221> misc_feature <222> (1444)..(1444) <223> "n" is any nucleotide <400> 197 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgcat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180 getaggeteg tegtettgge caeggecaec ceteceggea gtgtgacaac geeceaecee 240

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Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Thr Gly Asp Val Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160

Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190 Glu Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Trp
195 200 205

Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220

Gly Glu Arg Pro Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240

Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255

Val Arg Leu Arg Ala Tyr Xaa Asn Thr Pro Gly Leu Pro Val Cys Gln 260 265 270

Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285

Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Gly Glu Asn Phe Pro 290 295 300

Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315

Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335

Xaa Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350

Asn Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365

Met Ser Ala Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly 370 375 380

Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val 385 390 395 400

Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415

Asp Arg Glu Val Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser

420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 440 435 Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 455 Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 475 465 470 Trp Xaa Thr Tyr <210> 199 <211> 1485 <212> DNA <213> hepatitis C virus <220> <221> misc_feature <222> (684)..(684) <223> "n" is any nucleotide <220> <221> misc_feature <222> (909)..(909) <223> "n" is any nucleotide <220> <221> misc_feature <222> (1058)..(1058) <223> "n" is any nucleotide <220> <221> misc_feature <222> (1133)..(1133) <223> "n" is any nucleotide <400> 199 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgtat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180

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Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Val 20 25 30

Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160 Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190

Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg 195 200 205

Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220

Gly Glu Arg Xaa Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240

Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr
245 250 255

Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln
260 265 270

Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285

Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Xaa Pro 290 295 300

Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320

Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335

Thr Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350

Xaa Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365

Met Ser Ala Asp Leu Glu Val Ile Thr Xaa Thr Trp Val Leu Val Gly 370 375 380

Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val

205	390	205	400
385	370	395	400

Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415

Asp Arg Glu Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser 420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe
435 440 445

Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 450 455 460

Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 465 470 475 480

Trp Ala Thr Tyr

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<211> 340

<212> DNA

<213> hepatitis C virus

<400> 201

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<210> 202

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 202

Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Val Tyr Gln
1 5 10 15

360

420

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 120 115 Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp Gly Ile Asn Tyr Ala Thr <210> 165 499 <211> <212> DNA hepatitis C virus <213> <220> <221> misc_feature (306) . . (306) <222> <223> "n" is any nucleotide <220> misc feature <221> (313)..(313) <222> "n" is any nucleotide <223> <220> <221> misc_feature (340)..(340) <222> "n" is any nucleotide <223> atgagcacga atcctaaacc tcaaagaaaa accaaacgta acaccaaccg ccgccctatg 60 120 gacgttaagt tcccaggcgg tggtcagatc gttggcggag tttacttgtt gccgcgcagg ggccccaggt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcgtggg 180 aggegecaae etateeecaa ggegegeega acegagggea gateetggge geageeeggg 240 300 tatccttggc ccctttacgg caatgagggc tgtgggtggg cagggtggct cctgtcccct

cgcggntctc ggncgtcttg gggccccaat gatccccggn ggagatcccg caacttgggt

aaggtcatcg ataccctaac atgcggcttc gccgacctca tgggatacat cccgcttgta

ggcgcccccg tgggtggcgt cgccagggcc ctggcacatg gtgttagggc tgtggaagac

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499

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<211> 126

<212> PRT

<213> hepatitis C virus

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<221> MISC FEATURE

<222> (102)..(102)

<223> "Xaa" is any amino acid

<220>

<221> MISC FEATURE

<222> (105)..(105)

<223> "Xaa" is any amino acid

<220>

<221> MISC FEATURE

<222> (114)..(114)

<223> "Xaa" is any amino acid

<400> 166

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Thr Glu Gly Arg Ser Trp Ala Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Xaa Ser Arg Xaa Ser Trp Gly Pro Asn Asp Pro 100 105 110



Arg Xaa Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu
115 120 125

<210> 167 <211> 579 <212> DNA <213> hepatitis C virus <400> 167

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<210> 168 <211> 193 <212> PRT

<213> hepatitis C virus

<400> 168

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr
130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<210> 169

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 169

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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 170

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1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Gly Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Asp Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Val Gly Ala Pro 115 120 . 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Arg Met 180 185 190 <210>

<211><212>

171

579

DNA

hepatitis C virus

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Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Leu Thr Asn Asp

60

120

180

240

300

360

420

480

540

65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Val His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro Tyr Leu Gly Ala Pro
115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr
130 135 140

Ala Gly Gln Met Phe Thr Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Thr Tyr Pro Gly His Ile Thr Gly His Arg Met 180 185 190

Ala ·

<210> 173

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 173

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579

tccatctatc caggccacat cacgggtcac agaatggcg

<210> 174

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 174

Thr Cys Gly Ser Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Glu His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Ile Ala Leu Thr Pro Thr Leu Ala Ala Pro His Ile Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Ile Phe Leu 145 150 155 160

Val Gly Gln Met Phe Asn Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met

180 185 190

Ala

<210> <211>	1/5 579						
	DNA						
<213>	hepa	atitis C vi	cus				
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gggaato	ttc	ccggttgctc	cttttctatc	ttccttctag	cacttctctc	gtgcttgact	180
gtcccgg	gcct	cggcgcagca	ctaccggaac	atctcgggca	tttatcacgt	caccaatgac	240
tgcccga	act	ctagtatagt	gtatgaagct	gaccatcata	tcatgcatct	accagggtgt	300
gtgcctt	gcg	tgagaaccgg	gaacacctcg	cgctgctggg	ttcctttaac	acccactgtg	360
gctgccc	cct	atgttggcgc	gccgctcgaa	tccatgcggc	ggcacgtgga	cttaatggtg	420
ggtgccg	gcca	ccgtctgctc	ggccctgtac	atcggagacc	tttgcggagg	tgtcttcctg	480
			ccggccgcgc				540
greggge	aga	cgcccacccc	ccggccgcgc	cyccaccyya	0000000	00500000	• • •
tctatct	atg	atggccacat	caccggccat	agaatggct			579
<210>	176						

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 176

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys 100 105 Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 150 155 145 Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg Met Ala <210> 177 <211> 579 <212> DNA <213> hepatitis C virus <400> 177 acgtgcgggt tcgccgacct catgggatac atcccgctcg tgggcgctcc agtaggaggc 60 gtcgccagag ccttggcgca tggcgtcagg gctgtggagg acgggatcaa ttacgcaaca gggaacette ceggetgete ettitetate tiectetigg tactietete gegeetaact 180 gtcccagcgt ctgctcagca ctaccggaat gcatcgggca tctaccatgt caccaacgac 240 tgcccgaact ccagtattgt gtatgaagcc gaccatcaca tcatgcacct acccgggtgt 300

gtgccctgtg taagaactgg gaatgtctcg cgttgctgga ttcctttaac acccactgta

geogteecet acetegggge tecaettacg tetgtacgge ageatgtgga cetgatggtg

ggggcggcca ccttatgctc tgccctctac atcggagacc attgcggagg tgtcttcttg

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp

75

70 .

360

420

qcaqqqcaga tggtcagttt ccaaccccgg cgtcattgga ctacccagga ttgcaactgt tccatctatg tgggccacat caccggccac aggatggcc 579

<210> 178

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 178

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 10

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 45 35

Ser Ile Phe Leu Leu Val Leu Leu Ser Arg Leu Thr Val Pro Ala Ser 55 . 60

Ala Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 75

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His . 90 85

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Val Ser Arg Cys 105 110 100

Trp Ile Pro Leu Thr Pro Thr Val Ala Val Pro Tyr Leu Gly Ala Pro 120 115

Leu Thr Ser Val Arg Gln His Val Asp Leu Met Val Gly Ala Ala Thr 130 135

Leu Cys Ser Ala Leu Tyr Ile Gly Asp His Cys Gly Gly Val Phe Leu

Ala Gly Gln Met Val Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165

Asp Cys Asn Cys Ser Ile Tyr Val Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

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<210> 179
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<212> DNA
<213> hepatitis C virus
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<221> misc_feature
<222> (568)..(579)
<223> "n" is any nucleotide
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                                                                     120
qqqaatcttc ctggttgctc tttctctatc tcccttcttg aacttctctc gtgcctgact
                                                                     180
gttcccgcct cagccatcca ctatcgcaat gcttcggacg gttattatat caccaatgat
                                                                     240
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                                                                     300
ataccetgtg tgaagaccgg gaatcagteg eggtgetggg tggeteteac ecceaegetg
                                                                     360
geggeeecac acetaegtge teegettteg teettaeggg egeatgtgga eetaatggtg
                                                                     420
ggggccgcca cggcatgctc cgctttttac attggagatc tgtgcggggg tgtgtttttg
                                                                     480
gegggecaac tgttcactat eeggecaege atteatgaaa eeacteagga etgeaattge
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tccatctact cagggcacat cacgggtnnn nnnnnnnn
                                                                     579
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<211> 193
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<213> hepatitis C virus
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<223> "Xaa" is any amino acid
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<221> MISC FEATURE
<222> (190)..(193)
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<223> "Xaa" is any amino acid

<400> 180

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
1 10 15

Pro Val Gly Gly Val Ala Arg Xaa Leu Ala His Gly Val Arg Ala Leu 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Ser Leu Leu Glu Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Ile His Tyr Arg Asn Ala Ser Asp Gly Tyr Tyr Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Glu Asn His Ile Leu His
85 90 95

Leu Pro Gly Cys Ile Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro His Leu Arg Ala Pro 115 120 125

Leu Ser Ser Leu Arg Ala His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Ala Cys Ser Ala Phe Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Leu Phe Thr Ile Arg Pro Arg Ile His Glu Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly Xaa Xaa Xaa 180 185 190

Xaa

<210> 181 <211> 579

<212> DNA <213> hepatitis C virus <220> <221> misc_feature (152)..(152) <222> "n" is any nucleotide <223> <400> 181 gcgtgcggct tcgccgatct catgggatac atcccgctcg taggcgcccc cgtgggtggc 60 gtcgccagag ccctggcgca cggtgttagg gctgtggagg acgggattaa ctacgcaaca 120 180 gggaatette etggttgete tttetetate tneettetgg eacttetete gtgeetgaet 240 qtcccqqcct cggctcagca ctaccggaat gtctcgggca tctaccacgt caccaatgat 300 tgcccqaatt ccagcatagt gtatgaagcc gatcaccaca tcatgcactt accagggtgc ataccetgeg tgaggacegg gaacgttteg egetgetggg tatetetgae acetaetgtg 360 gctgctccct accteggggc tecgettacg tegetacgge ggcatgtgga tttgatggtg 420 qqtqcagcca ccctttgctc tgccctctac gtcggagacc tctgtggagg tgtcttccta 480 gtgggacaga tgttcacctt ccagccgcgc cgccactgga ccactcagga ctgcaactgc 540 579 tccatttacg tcggccacat cacaggccac agaatggct <210> 182 193 <211> PRT <212> <213> hepatitis C virus <220> <221> MISC FEATURE (51)..(51) <222> "Xaa" is any amino acid <223>

<400> 182

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Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Xaa Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser

B 7 -	~ 1~	uic	Tree	Ara	Asn	Va 1	Sar	Glv	Tla	ጥረታ	Hic	Val	Thr	Δen	Asn		
65	GIII	uis	ıyı	·	70	vai	Ser	GIY	116	75	1113	vai	1111	ASII	80		
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr		Ala 90	Asp	His	His	Ile	Met 95	His		
Leu	Pro	Gly	Cys 100	Ile	Pro	Cys	Val	Arg 105		Gly	Asn	Val	Ser 110	Arg	Cys		
Trp	Val	Ser 115	Leu	Thr	Pro	Thr	Val 120	Ala	Ala	Pro	Tyr	Leu 125	Gly	Ala	Pro		
Leu	Thr 130	Ser	Leu	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr		
Leu 145	Cys	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160		
Val	Gly	Gln	Met	Phe 165	Thr	Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln		
Asp	Cys	Asn	Cys 180	Ser	Ile	Tyr	Val	Gly 185	His	Ile	Thr	Gly	His 190	Arg	Met		
Ala																	
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gtc	gccag	ggg d	ccta	agaad	ca cg	gtgt	tagg	gct	gtgg	jagg	acgo	tatt	aa t	tate	caaca	*	120
ggga	atct	caa a	ccggt	tgct	c tt	tttc	ctato	tco	ctct	tgg	cact	tctt	tc g	gtgco	tgact		180
gtto	cca	cct o	cagco	cgtca	aa ct	atc	gcaac	gco	tcgg	gcg	tcta	tcat	at o	acca	atgac		240
tgc	cgaa	att o	gago	catag	gt gt	acga	aggct	gad	taco	aca	tcct	acac	ct o	ccts	ggtgc		300

ttaccctgcg tgagggttgg gaatcagtca cgctgctggg tggcccttac tcccaccgtg 360

geggegeett aegttggtge teegetagaa teeeteegga gteatgtgga tetgatggta qqtqctgcta ctgtgtgctc cgctctttac atcggggacc tgtgcggtgg cgtatttttg gttggtcaga tgttttcttt ccagccgcga cgccactgga ccacgcagga ctgcaattgt tctatctacg cggggcacgt tacgggccac aggatggca <210> 184 <211> 193 <212> PRT <213> hepatitis C virus <400> 184 Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 70 75 65 Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 90 Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 135 130 Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 160 150 145

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln

420

480

540

165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180

Ala

<211> 57	85 79				•		
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gcggcgcct	tt aca	tcggtgc	tccagttgaa	tccttccgga	gacacgtgga	catgatggtg	420
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gttggtcag	ga tgt	tttcttt	ccggccacga	cgccactgga	ctacgcagga	ctgcaattgt	540
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<210> 186

<211> 193 <212> PRT

<213> hepatitis C virus

<400> 186

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 5 10

Pro Val Gly Gly Val Ala Arg Ala Leu Glu His Gly Val Arg Ala Val 20

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 40

Ser Ile Tyr Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser • 55 Ala Ile His Tyr Arg Asn Ala Ser Gly Val Tyr His Val Thr Asn Asp 70 Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Leu His 90 85 Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys 105 Trp Val Ala Leu Ser Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 120 Val Glu Ser Phe Arg Arg His Val Asp Met Met Val Gly Ala Ala Thr 135 Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 150 145 Val Gly Gln Met Phe Ser Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Gly Met 180 Ala <210> 187 <211> 579 <212> DNA <213> hepatitis C virus <400> 187 acttgcggct ttgccgacct catgggatac atcccgctcg taggcgcccc tgtgggtggc 60 gtcgccaggg ccctggcaca cggtgttagg gctgtggagg acgggatcaa ttatgcgaca 120 gggaatette eeggttgete titetetate tieetetigg eactietite gigeetgaet 180 gttcccacct cggccgtcaa ctatcgcaat gcctcgggca tctatcacat caccaatgac 240 tgcccgaact cgagcatagt gtacgagacc gagcaccaca tcctacacct cccagggtgt 300 ttaccctgcg tgagggttgg gaatcagtca cgctgctggg tggccctcac tcccaccgtg 360

geggegeett acateggege teegettgaa teeeteegga gteatgtgga tetgatggta ggtgccgcta ctgcgtgctc cgctctttac atcggagacc tgtgcggtgg cgtatttttg gttggtcaga tgttctcttt ccagccgcgg cgccactgga ctacgcagga ctgcaattgt tccatctacg cggggcacgt tacgggccac aggatggca <210> 188 <211> 193 <212> PRT <213> hepatitis C virus <400> 188 Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 10 5 Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 55 Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 70 75 Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His 85 Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys 105 100 110 Trp Val Ala Leu Thr Pro Thr Val Ala Pro Tyr Ile Gly Ala Pro 125 115 120 Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu

155

145

150

420

480

540

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210> <211> <212> <213>	189 579 DNA hepa	atitis C vi	rus	4.			
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gtcccas	gcgt	cagctgagca	ctaccggaat	gcttcgggca	tctatcacat	caccaatgac	240
tgtccga	aatt	ccagcgtagt	ctatgaaact	gaccaccata	tattgcactt	gccggggtgc	300
gtaccct	gcg	tgagggccgg	gaacgtgtct	cgttgctgga	cgccggtaac	acctacggtg	360
gctgccg	gtat	ccatggacgc	teegetegag	teetteegge	ggcatgtgga	cctaatggta	420
ggtgcgg	gcca	ccgtgtgttc	tgtcctctat	gttggagacc	tctgtggagg	tgctttccta	480
gtgggg	caga	tgttcacctt	ccagccgcgt	cgccactgga	ccacgcagga	ttgtaattgc	540
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<210> 190

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 190

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Ala	Ser			
Ala 65	Glu	His	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thr	Asn	Asp 80			
Cys	Pro	Asn	Ser	Ser 85	Val	Val	Tyr	Glu	Thr 90	Asp	His	His	Ile	Leu 95	His			
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Ala	Gly	Asn	Val	Ser 110	Arg	Cys			
Trp	Thr	Pro 115	Val	Thr	Pro	Thr	Val 120	Ala	Ala	Val	Ser	Met 125	Asp	Ala	Pro			
Leu	Glu 130	Ser	Phe	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr			
Val 145	Cys	Ser	Val	Leu	Туг 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Ala	Phe	Leu 160			
Val	Gly	Gln	Met	Phe 165	Thr	Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln			
Asp	Cys	Asn	Cys 180		Ile	Tyr	Thr	Gly 185	His	Ile	Thr	Gly	His 190	Arg	Met		•	
Ala																		
	-		titi	s C	viru	s												
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															cgcagg	120		
ggo	ссса	ıggt	tggg	jtgtg	ica c	gcga	ctag	g aa	gact	tcgg	agc	ggtc	gca	acct	cgtggg	180		
aga	acgto	agc	ctat	cccc	aa g	gcac	gtcg	a to	tgag	ggaa	ggt	cctg	ggc	tcag	cccggg	240		
tad	cccat	ggc	ctct	ttac	gg t	aatg	aggg	ıt tg	tggg	tggg	cag	gatg	199			289		

192 <210> 96 <211> PRT <212> hepatitis C virus <213> 192 <400> Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 40 Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 55 Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro Gly 70 75 Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp <210> 193 <211> 498 <212> DNA hepatitis C virus <213> <400> 193 60 atgagcacga atcctaaacc tcaaagaaaa accaaacgta acaccaaccg ccgccctatg qacqtaaaqt tcccgggcgg tggacagatc gttggcggag tttacttgtt gccgcgcagg 120 180 qqccccqqt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcgtggc 240 aggegteaac ctatececaa ggegegeegg teegagggea ggteetggge geaageeggg 300 tacccctggc ccctctatgg caatgagggc tgtgggtggg cagggtggct cctgtctcct 360 egeggetete ggecatettg gggeccaaat gateecegge ggagategeg caatetgggt 420 aaggtcatcg ataccctgac gtgcggcttc gccgacctca tgggatacat cccgctcgtg

ggcgccccg tcgggggcgt cgccagggcc ctggcgcatg gcgtcagggc tgtggaggac

gggattaact atcgacag

480

498

<210> 194

<211> 166

<212> PRT

<213> hepatitis C virus

<400> 194

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly
20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Ala Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp 145 150 155 160

Gly Ile Asn Tyr Arg Gln 165

<210> 195

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 195

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gggaatctgc	ctggttgctc	tttctctatc	ttcattcttg	cacttctctc	gtgcctcact	180
gtcccggcct	ctgcagttcc	ctaccgaaat	gcctctggga	tctatcatgt	caccaatgat	240
tgcccaaact	cttccatagt	ctatgaggca	gatgatctga	tectacacge	acctggctgc	300
gtgccttgtg	tcaggaaaga	taatgtgagt	aggtgctggg	tccaaattac	ccccacgctg	360
tcagccccga	gcttcggagc	agtcacggct	ccccttcgga	gagccgttga	ttacttggtg	420
ggagggctg	ccctctgctc	cgcgttatac	gttggagacg	cgtgtggggc	actatttttg	480
gtaggccaaa	tgttcaccta	taggcctcgc	cagcatgcta	cggtgcagga	ctgcaactgt	540
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<210> 196

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 196

Thr Cys Gly Phe Ala Asp Leu Val Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu
20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val

, , , , , , , , , , , , , , , , , , ,	
Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr 20 25 30	
Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45	
Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60	
Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80	· · · · · · · · · · · · · · · · · · ·
Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95	
Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Lys Arg Ala Leu Gly 100 105 110	
Ala	
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ggagcctgaa acccgcaagg taatatctgc cctcactgaa agactctatg tgggcggtcc catgcacaac agcaggggag acctatgcgg gtaccgtaga tgccgcgcga gcggcgtata	180
caccacaage tregggaaca etetgaegtg etteetcaag gecacagegg ecaccaaage	240
cgctggccta aaggactgca ccatgttggt gtgtggtgac gacttagtcg ttatcgccga	300
aagcgatggt gtcgaagagg accgccgagc cctcggagct	340
<210> 204 <211> 113 <212> PRT <213> hepatitis C virus	
<400> 204	
Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15	

Cys Cys Asp Leu Glu Pro Glu Thr Arg Lys Val Ile Ser Ala Leu Thr Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Arg Gly Asp Leu Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 55 Gly Asn Thr Leu Thr Cys Phe Leu Lys Ala Thr Ala Ala Thr Lys Ala 65 Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Arg Arg Ala Leu Gly Ala <210> 205 <211> 340 DNA <212> hepatitis C virus <213> <400> 205 ctccacggtg accgaaaggg atatcaggac cgaggaagag atctaccagt gctgcgacct 60 ggagcccgaa gcccgcaagg tgatatccgc cctaacggaa agactctacg tgggcggtcc 120 catgtacaac tccaaggggg acctatgcgg gcaacggagg tgccgcgcaa gcggggtcta 180 caccaccage ttegggaaca etgtaacgtg ttateteaag geegttgegg etaetaggge cgcaggtctg aaaggttgca gcatgctggt ttgtggagac gacttagtcg tcatctgcga 300 340 gageggegge gtagaggagg atgeaagage ceteegagee

<210> 206

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 206

Ser Thr Val Thr Glu Arg Asp Ile Arg Thr Glu Glu Glu Ile Tyr Gln
1 5 10 15

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr 20 Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 40 35 Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Val Ala Ala Thr Arg Ala Ala Gly Leu Lys Gly Cys Ser Met Leu Val Cys Gly Asp Asp Leu Val 90 85 Val Ile Cys Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 105 Ala <210> 207 <211> 340 <212> DNA hepatitis C virus <213> <400> 207 ctccacggtg actgaaaggg acattagggt cgaggaagag atctaccagt gctgtgacct 60 ggagcccgag gcacgcaagg tgatatccgc tctcacagaa agactctaca agggcggccc 120 catgtataac agcaaggggg acctatgcgg gcttcggagg tgccgcgcaa gcggggtata 180 caccacaagc ttcgggaaca cggtgacatg ctaccttaaa gccacagcag ccaccagggc 240 tgcagggctg aaagattgca ctatgctggt atgcggtgac gacttagtcg ttattgccga 300 340 aagcggtggc gtggaggagg acgcccgagc cctccgagcc

<210> 208

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 208

Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Ile Tyr Gln

Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr

Glu Arg Leu Tyr Lys Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu

Cys Gly Leu Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val

Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg

Ala

<210> 209

<211> 340

<212> DNA

<213> hepatitis C virus

<220>

<221> misc_feature

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<223> "n" is any nucleotide

<220>

<221> misc_feature

<222> (23)..(23)

<223> "n" is any nucleotide

<220>

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<222> (65)..(65)

<223> "n" is any nucleotide

<220>

<221> misc_feature

<222> (72)..(72)

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<220>
<221> misc feature
<222> (277)..(277)
<223> "n" is any nucleotide
<220>
<221> misc feature
<222> (337)..(337)
<223> "n" is any nucleotide
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ggagnccgat gnccgcaagg tcatcaacgc cctcacagag agactctacg tgggcggccc
                                                                     120
tatgcacaac agcaagggag acctgtgtgg catccgtaga tgccgcgcga gcggcgttta
                                                                     180
caccacgage tteggaaaca egetgaettg etaceteaaa gecacagegg ecaccaggge
                                                                     240
cgcgggcttg aaggattgca ccatgctggt ctgcggngac gacctggttg tcattgctga
                                                                     300
                                                                     340
gagcattggc atagacgagg acaagcaagc cctccgnact
<210> 210
<211> 113
<212> PRT
<213> hepatitis C virus
<220>
<221> MISC_FEATURE
      (8)..(8)
<222>
<223> "Xaa" is any amino acid
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<221> MISC_FEATURE
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<223> "Xaa" is any amino acid
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<222>
<223> "Xaa" is any amino acid
<400>
Pro Thr Val Thr Glu Arg Asp Xaa Arg Val Glu Glu Val Tyr Gln
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10

Cys Cys Asn Leu Glu Xaa Asp Xaa Arg Lys Val Ile Asn Ala Leu Thr 20 25

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 40

Cys Gly Ile Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 55

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 70

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85

Val Ile Ala Glu Ser Ile Gly Ile Asp Glu Asp Lys Gln Ala Leu Arg 100

Thr

<210> 211

<211> 340

<212> DNA

<213> hepatitis C virus

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<223> "n" is any nucleotide

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<221> misc_feature

<222> (327)..(327)

<223> "n" is any nucleotide

<220>

<221> misc_feature

<222> (333)..(334)

<223> "n" is any nucleotide

<220>

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<222> (340)..(340)

<223> "n" is any nucleotide

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catqttcaac agcaagggag acctgtgcgg acaacgccgg tgccgcgcaa gcggcgtgtt
caccaccage ttegggaaca cactgaegtg ctacettaaa gecacagetg ctactagage
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gagcgccggt gtggaggagg atcccanaac ccnncgaccn
<210> 212
<211>
      113
<212> PRT
<213> hepatitis C virus
<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> "Xaa" is any amino acid
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<221> MISC_FEATURE
<222> (109)..(109)
<223> "Xaa" is any amino acid
<220>
<221> MISC FEATURE
<222> (111) ...(111)
<223> "Xaa" is any amino acid
<400> 212
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                                    10
Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr
                                25
            20
                                                    30
Glu Arg Leu Tyr Val Gly Gly Pro Met Phe Asn Ser Lys Gly Asp Leu
        35
                                                45
Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Phe
    50
                        55
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Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

60

120

180

240

300

340

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ser Glu Ser Ala Gly Val Glu Glu Asp Pro Xaa Thr Xaa Arg 100 105 110

Pro

<210>	213			
<211>	340			
<212>	DNA			
<213>	hepatitis	С	virus	

22135 Nepacicis C Viids

<400> 213
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ggcccccgag gccagacagg ccataaagtc gctcacagag cggctttata tcgggggtcc 120

cctgactaat tcaaaggggc agaactgtgg ctatcgccga tgccgcgcaa gcggcgtgct 180

gacgaccagc tgcggtaata cccttacatg ttacctaaag gcctctgcag cctgtcgagc 240

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aagcgcggga acccaagagg acgcggcgag cctacgagtc 340

<210> 214 <211> 113 <212> PRT <213> hepatitis C virus

<400> 214

Ser Thr Val Thr Glu Asn Asp Ile Arg Val Glu Glu Ser Ile Tyr Gln 1 5 10 15

Cys Cys Asp Leu Ala Pro Glu Ala Arg Gln Ala Ile Lys Ser Leu Thr 20 25 30

Glu Arg Leu Tyr Ile Gly Gly Pro Leu Thr Asn Ser Lys Gly Gln Asn 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Cys 50 55 60

Gly 65	Asn	Thr	Leu	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Cys	Arg	Ala 80	
Ala	Lys	Leu	Gln	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Cys	Glu 100	Ser	Ala	Gly	Thr	Gln 105	Glu	Asp	Ala	Ala	Ser 110	Leu	Arg	
Val																
<210 <210 <210 <210	1> 3 2> I	215 340 DNA nepat	titis	3 C T	/irus	·		. •								
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gcc	ccaa	gag g	gccag	gaact	g to	atac	cacto	c gct	caco	gag	agad	etcta	acg t	ggga	agggcc	120
cat	gataa	aac a	agcaa	aaggg	gc aa	tcct	gegg	g tta	acago	gcgt	tgco	gcgo	aa g	gcggt	gtttt	180
cac	cacca	agc a	atggg	ggaat	a co	atga	acgto	g tta	acato	caaa	gcco	ttgo	ag d	gtgt	aaagc	240
cgca	aggga	atc 9	gtgga	accc	g to	atgo	tggt	gtg	gtgga	gac	gaco	tggt	cg t	cato	tcgga	300
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Ala	Cys	Ser	Leu 20	Pro	Gln	Glu	Ala	Arg 25	Thr	Val	Ile	His	Ser 30	Leu	Thr	
Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Ile	Asn	Ser	Lys 45	Gly	Gln	Ser	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	

Gly 65	Asn	Thr	Met	Thr	Cys 70	Tyr	Ile	Lys	Ala	Leu 75	Ala	Ala	Cys	Lys	Ala 80	
Ala	Gly	Ile	Val	Asp 85	Pro	Val	Met	Leu	Val 90	Сув	Gly	Asp	Asp	Leu 95	Val	
Val	Ile	Ser	Glu 100	Ser	Gln	Gly	Asn	Glu 105	Glu	Asp	Glu	Arg	Asn 110	Leu	Arg	
Ala	·															
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tgaa	ccg	gag g	gccag	ggaaa	ag to	gatct	cct	c cct	cacc	gag	cggd	ettta	act g	gcgga	ggccc	120
tate	gttta	aac a	agcaa	a gggg	gg co	cagt	gtgg	g tta	atcgo	cgt	tgc	gtgo	ca g	gtgga	gttct	180
gcct	acca	agc t	tttgg	gcaad	a ca	atca	actt	g tta	acato	aag	gçca	acaac	gg (ccgcc	jaaggc	240
cgca	aggc	ctc	cggaa	acccg	gg ac	tttc	ttgt	ctg	gegga	gat	gato	tggt	.cg t	ggtg	gctga	300
gagt	gat	ggc g	gtcga	acgag	gg at	agaç	gcago	c cct	gaga	gcc						340
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Cys	Cys	Asn	Leu 20	Glu	Pro	Glu	Ala	Arg 25	Lys	Val	Ile		Ser 30	Leu	Thr	
Glu	Arg	Leu 35	Tyr	Cys	Gly	Gly	Pro 40	Met	Phe	Asn	Ser			Ala	Gln	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Leu 60	Pro	Thr	Ser	Phe	

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Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Thr Thr Ala Ala Lys Ala
                    70
                                        75
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Val Val Ala Glu Ser Asp Gly Val Asp Glu Asp Arg Ala Ala Leu Arg
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Arg Ser Glu Gly Arg Thr Ser Trp Ala Gln
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Arg Thr Glu Gly Arg Thr Ser Trp Ala Gln
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Phe Leu Ser Gln Thr Lys Gln Gln Gly Leu Asn Phe Ser Phe Leu Thr 20 25 30

Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala Gln Ala Pro Pro Pro Ser 35 40 45

Trp Asp Glu Met Trp Lys Cys Leu Val Arg Leu Lys Pro Thr Leu His 50 55 60

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 65 70 75 80

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 85 90 95

Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu Gly Gly Val Leu 100 105 110

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Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys Val Val Ile Val
                            120
        115
Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val Pro Asp Lys Glu
    130
                       135
Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala
                   150
                                       155
Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys
                                    170
Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Gln Ala Val Ile Glu
           180
Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala Phe Trp His Lys
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Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val
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Gln His Tyr Arg Asn Val Ser Gly Ile Tyr His Val
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Val Tyr Glu Thr Asp Asn His Ile Leu His Leu 5

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Val Tyr Glu Thr Glu Asn His Ile Leu His Leu

<210> 236

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<400> 236

Val Phe Glu Thr Val His His Ile Leu His Leu

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Val Phe Glu Thr Glu His His Ile Leu His Leu
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Val Phe Glu Thr Asp His His Ile Met His Leu
1 5
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Val Tyr Glu Thr Glu Asn His Ile Leu His Leu
1 5
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Val Tyr Glu Ala Asp Ala Leu Ile Leu His Ala
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Ala Pro Ser Phe Gly Ala Val Thr Ala Pro
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Val Ser Gln Pro Gly Ala Leu Thr Lys Gly
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Ala Pro Tyr Ile Gly Ala Pro Val Glu Ser
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Ala Gln His Leu Asn Ala Pro Leu Glu Ser
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Ser Pro Tyr Val Gly Ala Pro Leu Glu Pro
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Ser Pro Tyr Ala Gly Ala Pro Leu Glu Pro
1 5
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Ala Pro Tyr Leu Gly Ala Pro Leu Glu Ser
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Ala Pro Tyr Val Gly Ala Pro Leu Glu Ser
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Ala Pro Tyr Leu Gly Ala Pro Leu Thr Ser
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Ser Pro Gln His His Lys Phe Val Gln Asp
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Pro Pro Arg Ile His Glu Thr Thr Gln Asp
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Thr Ile Ser Tyr Ala Asn Gly Ser Gly Pro Ser Asp Asp Lys
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<210> 268

hepatitis C virus

268 <400>

Ser Arg Arg Gln Pro Ile Pro Arg Ala Arg Arg Thr Glu Gly Arg Ser

Trp Ala Gln

269 <210>

<211> 1443

<212> DNA

hepatitis C virus

<400> 269

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Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr 35 40 45

Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala 50 55 60

Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu 65 70 75 80

Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Gly Arg Ala 85 90 95

Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His
100 105 110

Ser Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr Ser Leu Gly 115 120 125

Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala Val Ile Pro 130 135 140 Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala Gly Glu Arg Pro Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Phe Pro Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro Pro Pro Ser

Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro Thr Leu Thr 325 330 335

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 340 345 350

Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 355 360 365

Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly Gly Val Val 370 380

Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val Ala Ile Val 385 390 395 400

Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu 405 410 415

Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser Ala Ser Leu 420 425 430

Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe Lys Glu Lys
435
440
445

Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu Thr Leu Lys 450 455 460

Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe Trp Ala Thr 465 470 475 480

Tyr

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<211> 5

<212> PRT

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<400> 271

Leu Glu Trp Arg Asn 5

<210> 272

<211> 20

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Glu Phe Asp Glu

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 Phe Lys Gln Lys
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  Ser Arg Gln Ala
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    Glu Phe Asp Glu
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Phe Lys Gln Lys
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Thr Lys Gln Ala
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Ala Phe Asp Glu
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Ala Ser Arg Ala Ala Leu Ile Glu Glu Gly Gln Arg Ile Ala Glu Met
Leu Lys Ser Lys
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Ser Lys Gln Ala 20

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<213> hepatitis C virus

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Ala Phe Asp Glu 20

<210> 282

<211> 20

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Ala Ser Lys Ala Ala Leu Ile Glu Glu Gly Gln Arg Met Ala Glu Met 1 5 10 15

Leu Lys Ser Lys 20

<210> 283

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Met Ala Glu Met Leu Lys Ser Lys Ile Gln Gly Leu Leu Gln Gln Ala 1 5 10 15

Thr Arg Gln Ala 20

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<211> 20

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Phe Lys Glu Lys